

Supporting Information

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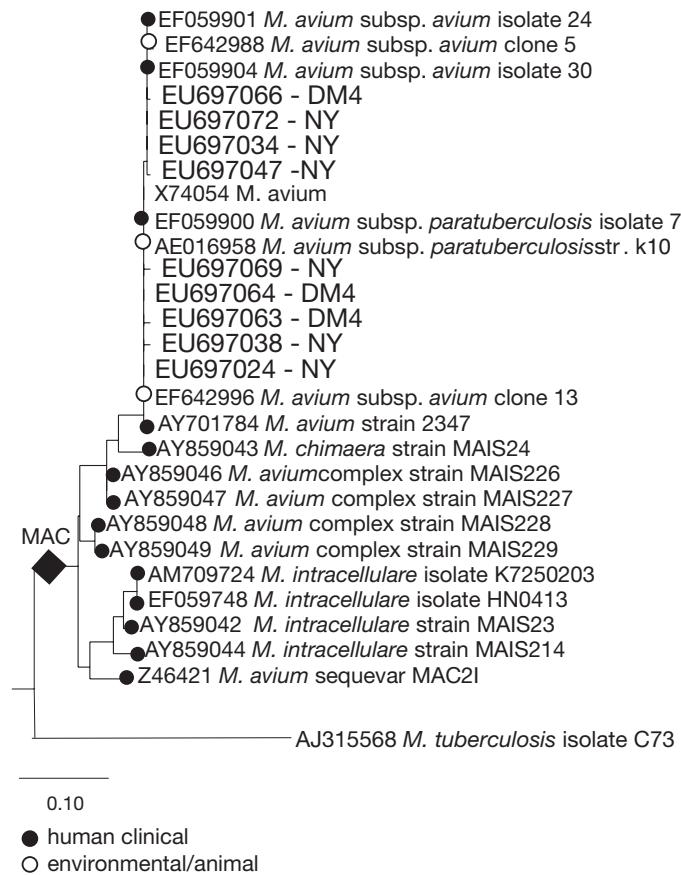


Fig. S1. Results of a phylogenetic analysis of the *M. avium* complex (MAC) using 16S-23S Internally Transcribed Spacer (ITS) gene sequences. Alignment includes cultured representatives of MAC from clinical and non-clinical sources (NCBI accession numbers are listed) as well as showerhead biofilm sequences from this study. Sequences from this study are denoted in bold and city of origin is indicated: DM4 (Denver Metro 4) and NY (New York). Tree does not show bootstrap values due to the statistically insignificant number of differences in the ITS sequences.

Other Supporting Information Files

- [Dataset S1](#)
- [Dataset S2](#)
- [Dataset S3](#)
- [Dataset S4](#)
- [Dataset S5](#)